

Whole-genome sequencing and antimicrobial resistance in *Brucella melitensis* from a Norwegian perspective.

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Supplementary information

Supplementary information. Calculated average coverage depth and coverage breadth for *Brucella melitensis* isolates. Calculated using SAMtools.

Isolate no.	Average coverage depth	Coverage breadth (% covered by 5 or more reads)
1	40.10	99.91%
2	82.05	99.93%
4	19.66	99.88%
5	71.94	99.91%
6	36.75	99.91%
7	45.65	99.91%
8	62.25	99.92%
9	65.79	99.95%
11	56.44	99.95%
13	32.29	99.90%
20	25.10	99.88%
23	76.91	99.92%
24	64.83	99.95%
36	39.36	99.91%
37	20.25	99.83%
40	44.69	99.92%
41	83.69	99.92%
43	110.18	99.91%
44	48.81	99.91%
45	25.29	99.86%
46	48.33	99.90%
47	18.06	77.99%
48	70.25	99.92%